

Sequence Listing

<110> OncoMab GmbH
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MULLER-HERMELINK, Hans-Konrad

<120> Human monoclonal antibody

<130> 043043-0359295

<140> 10/579,290

<141> 2004-11-12

<150> EP - 03 026 161.4

<151> 2003-11-14

<160> 4

<170> PatentIn version 3.3

<210> 1

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<212> PRT

<213> Homo sapiens

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<221> misc_feature

<223> variable region of the light chain (VL) of antibody SAM-6

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Ser Tyr Val Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly
1 5 10 15

Gln Thr Ala Ser Ile Thr Cys Ser Gly Asp Lys Leu Gly Asp Lys
20 25 30

Tyr Ala Cys Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu
35 40 45

Val Ile Tyr Gln Asp Ser Lys Arg Pro Ser Gly Ile Pro Glu Arg
50 55 60

Phe Ser Gly Ser Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser
65 70 75

Gly Thr Gln Ala Met Asp Glu Ala Asp Tyr Tyr Cys Gln Ala Trp
80 85 90

Asp Ser Ser Ile Val Val
95

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 <212> DNA
 <213> Homo sapiens

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 Ser Tyr Val Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly
 1 5 10 15

cag aca gcc agc atc acc tgc tct gga gat aaa ttg ggg gat aaa 90
 Gln Thr Ala Ser Ile Thr Cys Ser Gly Asp Lys Leu Gly Asp Lys
 20 25 30

tat gct tgc tgg tat cag cag aag cca ggc cag tcc cct gtg ctg 135
 Tyr Ala Cys Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Val Leu
 35 40 45

gtc atc tat caa gat agc aag cgg ccc tca ggg atc cct gag cga 180
 Val Ile Tyr Gln Asp Ser Lys Arg Pro Ser Gly Ile Pro Glu Arg
 50 55 60

ttc tct ggc tcc aac tct ggg aac aca gcc act ctg acc atc agc 225
 Phe Ser Gly Ser Asn Ser Gly Asn Thr Ala Thr Leu Thr Ile Ser
 65 70 75

ggg acc cag gct atg gat gag gct gac tat tac tgt cag gcg tgg 270
 Gly Thr Gln Ala Met Asp Glu Ala Asp Tyr Tyr Cys Gln Ala Trp
 80 85 90

gac agc agc att gtg gta 288
 Asp Ser Ser Ile Val Val
 95

<210> 3
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<220>
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Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro Gly
 1 5 10 15

Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser
 20 25 30

Ser Tyr Ala Met His Trp Val Arg Glu Ala Pro Gly Lys Gly Leu
 35 40 45

Glu Trp Val Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr
 50 55 60

Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser
65 70 75
Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp
80 85 90
Thr Ala Val Tyr Tyr Cys Ala Arg Asp Arg Leu Ala Val Ala Gly
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Lys Thr Phe Asp Tyr
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<210> 4
<211> 330
<212> DNA
<213> Homo sapiens
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Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro Gly
1 5 10 15
agg tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc agt 90
Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser
20 25 30
agc tat gct atg cac tgg gtc cgc cag gct cca ggc aag ggg ctg 135
Ser Tyr Ala Met His Trp Val Arg Glu Ala Pro Gly Lys Gly Leu
35 40 45
gag tgg gtg gca gtt ata tca tat gat gga agc aat aaa tac tac 180
Glu Trp Val Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr
50 55 60
gca gac tcc gtg aag ggc cga ttc acc atc tcc aga gac aat tcc 225
Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser
65 70 75
aag aac acg ctg tat ctg caa atg aac agc ctg aga gct gag gac 270
Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp
80 85 90
acg gct gtg tat tac tgt gcg aga gat cgg tta gca gtg gct ggt 315
Thr Ala Val Tyr Tyr Cys Ala Arg Asp Arg Leu Ala Val Ala Gly
95 100 105
aaa act ttt gac tac
Lys Thr Phe Asp Tyr
110 330